



Prevalence of β^S -globin gene haplotypes, α -thalassemia (3.7 kb deletion) and redox status in patients with sickle cell anemia in the state of Paraná, Brazil

Eliana LitsukoTomimatsu Shimauti^{1,2}, Danilo Grunig Humberto Silva³, Eniuce Menezes de Souza⁴, Eduardo Alves de Almeida³, Francismar Prestes Leal⁵ and Claudia Regina Bonini-Domingos¹

¹Laboratório de Hemoglobinas e Genética das Doenças Hematológicas, Departamento de Biologia, Universidade Estadual Paulista “Júlio de Mesquita Filho”, São José do Rio Preto, SP, Brazil.

²Departamento de Análises Clínicas e Biomedicina, Universidade Estadual de Maringá, Maringá, PR, Brazil.

³Departamento de Química e Ciências Ambientais, Universidade Estadual Paulista “Júlio de Mesquita Filho”, São José do Rio Preto, SP, Brazil.

⁴Departamento de Estatística, Universidade Estadual de Maringá, Maringá, PR, Brazil.

⁵Hemocentro, Maringá, PR, Brazil.

Abstract

The aim of this study was to determine the frequency of beta S-globin gene (β^S globin) haplotypes and alpha thalassemia with 3.7 kb deletion ($-\alpha^{3.7kb}$ thalassemia) in the northwest region of Paraná state, and to investigate the oxidative and clinical-hematological profile of β^S globin carriers in this population. Of the 77 samples analyzed, 17 were Hb SS, 30 were Hb AS and 30 were Hb AA. The β^S globin haplotypes and $-\alpha^{3.7kb}$ thalassemia were identified using polymerase chain reaction. Trolox equivalent antioxidant capacity (TEAC) and lipid peroxidation (LPO) were assessed spectrophotometrically. Serum melatonin levels were determined using high-performance liquid chromatography coupled to coulometric electrochemical detection. The haplotype frequencies in the SS individuals were as follows: Bantu- 21 (62%), Benin - 11 (32%) and Atypical- 2 (6%). Bantu/Benin was the most frequent genotype. Of the 47 SS and AS individuals assessed, 17% (n = 8) had the $-\alpha^{3.7kb}$ mutation. Clinical manifestations, as well as serum melatonin, TEAC and LPO levels did not differ between Bantu/Bantu and Bantu/Benin individuals ($p > 0.05$). Both genotypes were associated with high LPO and TEAC levels and decreased melatonin concentration. These data suggest that the level of oxidative stress in patients with Bantu/Bantu and Bantu/Benin genotypes may overload the antioxidant capacity.

Keywords: antioxidants, hemoglobinopathies, melatonin, sickle cell disease, thalassemia.

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Introduction

Hemoglobin S (Hb S) is produced by the substitution of thymine by adenine (GAG to GTG) in the sixth codon of the β globin gene, which results in the production of valine rather than glutamic acid. In homozygous form (Hb SS), this mutation is known as sickle cell anemia (SCA), and in heterozygous form (Hb AS) as sickle cell trait (Frenette and Atweh, 2007). SCA is characterized by chronic inflammation and recurrent ischemic-reperfusion events that may lead to an excess of free radicals and oxidative stress (Souza, 2001; Kaul *et al.*, 2004; Dasgupta *et al.*, 2006). A number of studies suggest that oxidative stress plays an

important role in the pathophysiology of SCA (Hebbel *et al.*, 1988; Naoum, 2000; de Oliveira Filho *et al.*, 2013). Due to its greater tendency for self-oxidation and oxidant generation, Hb S leads to increased lipid peroxidation (LPO) and lysis in the sickle cell membrane. The intensity of LPO and the reduction in antioxidant defense both appear to be related to the clinical severity of SCA (Dasgupta *et al.*, 2006; Shimauti *et al.*, 2010).

Melatonin, a potential antioxidant and anti-inflammatory agent, has been implicated in a number of pathological processes because of its ability to detoxify reactive oxygen species (ROS) and nitrogen species (RNS), and to stimulate antioxidant enzymes (Reiter *et al.*, 2000; Cuzocrea and Reiter, 2002; Allegra *et al.*, 2003; Mayo *et al.*, 2005; Tan *et al.*, 2007; Shimauti *et al.*, 2010), thereby exerting a protective effect against oxidative damage in different biological systems. The trolox equivalent

antioxidant capacity (TEAC), which assesses the response to free radical attack (Re *et al.*, 1999), and the level of thiobarbituric acid reactive species (TBARS), a reliable indicator of membrane lipid peroxidation or oxidative stress (Block *et al.*, 2002), are some of the most important and widely used biomarkers to assess oxidative damage.

The clinical course of SCA is heterogeneous and the clinical expression of the condition appears to be influenced by genetic features such as alpha thalassemia, the haplotypes in the beta globin gene cluster and Hb F levels (Steinberg and Embury, 1986; Zago and Pinto, 2007; Silva and Gonçalves, 2010; Shimauti *et al.*, 2011). Alpha thalassemia can be caused by the deletion of one or both alpha globin genes (α_1 and α_2) (Harteveld and Higgs, 2010), and the most common cause of the condition in Brazil is the deletion of the $-\alpha^{3.7\text{kb}}$ gene as a result of homologous recombination between misaligned chromosomes (Sonati *et al.*, 1991; Wagner *et al.*, 2010). Polymorphisms in the β^S globin gene cluster have been associated with at least five haplotypes that are classified according to the African and Middle Eastern regions where the genes are most commonly found. The Benin haplotype is associated with West Africa, while Bantu is most commonly seen in Eastern and south-central Africa. The Senegal and Cameroon haplotypes are associated with Atlantic West Africa and the African West Coast, respectively, while the Arab-Indian haplotype is found in India and the East Arabian Peninsula (Chebloune *et al.*, 1988; Nagel and Ranney, 1990; Elion *et al.*, 1992; Lapoumeroulie *et al.*, 1992). The Senegal and Arab-Indian haplotypes are associated with increased Hb F levels ($> 15\%$) and less severe SCA. The Benin haplotype, on the other hand, is associated with intermediate Hb F levels (5% to 15%) and a severe clinical course, while individuals with the Bantu haplotype presenting lower Hb F levels ($< 5\%$) and a worse clinical evolution (Powars, 1991; Elion *et al.*, 1992; Powars and Hiti, 1993; Galiza Neto and Pitombeira, 2003).

The intense miscegenation of the Brazilian population has resulted in regional differences in ethnic composition that have led to a heterogeneous distribution of the Hb S gene (Wenning *et al.*, 2000; Chinelato-Fernandes and Bonini-Domingos, 2005; Seixas *et al.*, 2008; Silva Filho *et al.*, 2010). Paraná state, located in southern Brazil, has a 1.52% prevalence for the heterozygous S gene (Watanabe *et al.*, 2008). Hb S was introduced in Brazil during the African slave trade from the 16th to 19th centuries. Historical records state that most slaves from Benin went to the port of Salvador, in the state of Bahia, while Rio de Janeiro received slaves from the Angola region, where the Bantu haplotype was the most prevalent (Silva Filho *et al.*, 2010). After arriving in Rio de Janeiro, slaves were redistributed to other regions in the country, and studies show that most states in Brazil, especially those in the southeast and southern regions of the country, received slaves from Rio de Janeiro (Fleury, 2007). There are also records of slave trading

in the state of Paraná in the 17th and 18th centuries, and starting from the second half of the 18th century, internal migration of former slaves was observed from other regions of Brazil to Paraná (Luna and Klein, 2004). Paraná has a multiethnic population, a large percentage of which is of European origin. The percentage of Afro-Brazilians in Paraná was thought to be quite small, until a survey by the Brazilian Institute of Geography and Statistics revealed that these individuals made up 24% of the state's population (IBGE, 2013). These findings showed that Paraná is the southern Brazilian state with the largest proportion of Afro-Brazilian individuals (Gomes Júnior *et al.*, 2008).

The goal of the present study was to analyze the frequency of β^S gene haplotypes and alpha thalassemia ($-\alpha^{3.7\text{kb}}$) and to verify their association with oxidative stress biomarkers and the antioxidant defense system, in addition to establishing the clinical and hematological profile of β^S globin gene carriers in Paraná.

Subjects and Methods

Subjects

Seventy-seven individuals, regardless of gender, were randomly selected from the northwest region of the state of Paraná in southern Brazil. Of these participants, 17 were Hb SS carriers (aged 10 to 39 years) in a stable phase of the disease selected from outpatient clinics, 30 were carriers of Hb AS (14 to 53 years) selected from voluntary blood donors and 30 were healthy Hb AA carriers (11 to 55 years) who had total Hb levels within normal limits for their gender and age and were recruited as a control group (Lewis *et al.*, 2006). Eligible participants were non-smokers, not pregnant, non-alcoholics, in the stable phase of the disease, and had not received blood transfusions in the three months prior to participating in the study. Data regarding medication use, clinical events and blood transfusions were obtained by evaluating the subjects' medical records and by applying questionnaires at the time of blood collection. The study protocol was approved by the Research Ethics Committee of the Department of Life Sciences, Linguistics and Exact Science of the State University of São Paulo (UNESP), Brazil (protocol number 0025.0.229.000-07).

Biological samples

After informed consent, 20 mL of peripheral blood was collected and distributed into the following tubes: a) a tube with K3EDTA (4 mL) for hematological, chromatographic and molecular analyses, b) a tube with heparin (7 mL) for determination of plasma biomarkers of oxidative stress and antioxidant capacity, after centrifugation at 1500 rpm for 20 min, and c) a tube without anticoagulant (9 mL) to obtain serum for the quantification of melatonin. All biological samples were collected at 6:30 a.m. to pre-

vent the inhibition of melatonin secretion caused by exposure to daylight.

Hemoglobin profile and hematological parameters

Hematological parameters were obtained using an Auto Hematology Analyzer (BC-300 PLUS - Mindray, China) and microscopic analysis was done using blood smears stained according to the May-Grunwald-Giemsa method (Lewis *et al.*, 2006). The tests used to screen for hemoglobinopathy included hemoglobin electrophoresis at pH 8.4 (Marengo-Rowe, 1965) and pH 6.2 (Vella, 1968). The quantification of hemoglobin fractions was performed by high-performance liquid chromatography (HPLC) using a Variant chromatography system (Bio-Rad) (Instruction Manual, 2006).

Identification of Hb S genotypes, β^S gene haplotypes and $-\alpha^{3.7kb}$ thalassemia

The Hb S genotypes were confirmed by molecular analysis with polymerase chain reaction- restriction fragment length polymorphism (PCR-RFLP). The segment that encodes the β^S gene was amplified using specific primers and the amplified segment was cleaved with *DdeI* restriction endonuclease (New England Biolabs, MA, USA) (Bonini-Domingos, 2006). The β^S globin haplotypes were determined by PCR-RFLP with an assessment of six polymorphic restriction sites: 5' γ^G (*XmnI*), γ^G (*HindIII*), γ^A (*HindIII*), $\psi\beta$ (*HincII*), 3' $\psi\beta$ (*HincII*) and 5' β (*HinfI*), according to the method reported by Sutton *et al.* (1989). The $-\alpha^{3.7}$ deletion was identified by multiplex PCR as described by Chong *et al.* (2000).

Biochemical analysis

Lipid peroxidation was assessed based on the plasma TBARS levels, according to the method described by Mihara and Uchiyama (1978). Plasma antioxidant capacity was determined by the TEAC method, which involves the use of Trolox (6-hydroxy-2,5,7,8-tetramethylchroman-2-carboxylic acid, product no. 23881-3; Aldrich Chemical Co.), a synthetic water-soluble antioxidant analogous to vitamin E (Miller *et al.*, 1993; Re *et al.*, 1999). The serum melatonin concentration was assessed using HPLC coupled to a coulometric electrochemical detector (Coulochem III ESA model 526, Bedford, MA, USA), as previously reported (Shimauti *et al.*, 2010).

Statistical analysis

Statistical analyses were done using Statistica software, version 8.0. The data were assessed for normality and homoscedasticity, and were analyzed using the non-parametric Mann-Whitney and Kruskal-Wallis tests, complemented by Dunn's test. Correlation analyses were done using Spearman's test. The significance level for the tests was set at 5% ($\alpha = 0.05$).

Results

Nine (52.9%) individuals with SCA (Hb SS) used hydroxyurea (HU). However, as there were no significant differences in Hb F levels ($p = 0.199$), TEAC ($p = 0.832$), melatonin ($p = 0.962$) and TBARS ($p = 0.835$) between HU users and nonusers, these individuals were combined into a single group. Analyses also showed that the age and gender of the individuals in the Hb SS, Hb AS and Hb AA groups did not influence the levels of redox status biomarkers ($p > 0.05$).

Alpha thalassemia was analyzed in 47 individuals (17 Hb SS and 30 Hb AS), eight of whom (17%) had the $-\alpha^{3.7kb}$ mutation. Seven (14.9%) individuals were heterozygotes ($-\alpha/\alpha$) and one (2.1%) was homozygous ($-\alpha/-\alpha$) for the mutation (Table 1). The β^S gene polymorphisms were analyzed in 17 Hb SS patients. The Bantu haplotype was the most frequent in these individuals and was found in 21 (62%) chromosomes, while the Benin haplotype was found in 11 (32%) chromosomes (Table 2). The Senegal and Arab-Indian haplotypes were not detected in the sample.

The subjects were divided into Bantu/Bantu and Bantu/Benin groups in order to analyze the influence of haplotypes on the oxidative and hematologic parameters and on phenotypic expression. Table 3 shows the redox profile and hematologic characteristics of the entire sample. The serum melatonin concentrations, TEAC and LPO in Hb AS individuals were not statistically different from those in subjects with Hb AA genotypes ($p > 0.05$). However, a marked reduction in serum melatonin levels ($p < 0.001$) and an elevation in TEAC ($p \leq 0.007$) and LPO ($p < 0.001$) were observed in Hb SS subjects as compared to Hb AS and Hb AA individuals. The antioxidant/oxidant and hematologic profiles did not differ significantly between the Bantu/Bantu and Bantu/Benin groups ($p > 0.05$), except for the mono-

Table 1 - Frequency of the $-\alpha^{3.7kb}$ thalassemia mutation in β^S -globin gene carriers.

Genotypes	$\alpha\alpha/\alpha\alpha$ N (%)	$-\alpha/\alpha\alpha$ N (%)	$-\alpha/-\alpha$ N (%)	N
SS	14 (82.4)	2 (11.8)	1 (5.8)	17
AS	25 (83.3)	5 (16.7)	0	30
Total	39 (83.0)	7 (14.9)	1 (2.1)	47

Table 2 - Frequency of β^S chromosomes and genotypes in 17 Hb SS individuals.

Chromosomes	N (%)	Haplotypes	N (%)
Bantu	21 (62)	Bantu/Bantu	6 (35.3)
Benin	11 (32)	Bantu/Benin	7 (41.1)
Atypical	2 (6)	Benin/Benin	2 (11.8)
		Bantu/Atypical	2 (11.8)
Total	34 (100)	Total	17 (100)

Table 3 - Demographic and laboratory features of individuals with sickle cell anemia (Hb SS), sickle cell trait (Hb AS) and without hemoglobinopathy (Hb AA).

Features	Hb SS (N = 17) Med (min-max)	Hb AS (N = 30) Med (min-max)	Hb AA (N = 30) Med (min-max)
Age	22(10-39)	37(14-53)	27(11-55)
TBARS (ng/mL)	863.4 ^a (496.8-2203)	470.6 ^b (280.8-2796.1)	485 ^b (270.2-676.3)
TEAC (mM)	2.00 ^a (1.86-2.10)	1.9 ^b (1.7 -2.0)	1.9 ^b (1.7-2.0)
Melatonin (pg/mL)	42.6 ^a (9.63-234.9)	116.2 ^b (6.63-633.8)	146 ^b (39.9-671.6)
Hb (g/dL)	7.6 ^a (5.9-9.7)	12.4 ^b (10.0-15.1)	13.7 ^b (11.5-15.4)
MCV (fL)	96.8 ^a (79.5-121.9)	84.8 ^b (76.0-94.0)	86.5 ^b (82-91.4)
MCH (pg)	31.2 ^a (24.8-40.4)	27.6 ^b (24.2-31)	28.5 ^b (27-30.1)
MCHC (%)	32.6 ^a (31-35.6)	32.8 ^a (31-34)	33.4 ^a (31.2-34.5)
Reticulocytes (%)	14.8 ± 8.8(6.3-36.9)	-	-
Leukocytes (x10 ⁹ /L)	12,600 ^a (8,600-18,800)	7,650 ^b (4,300-10,900)	6,550 ^b (3,800-10,500)
Neutrophils (x10 ⁹ /L)	6,090 ^a (4,472-11,280)	4,002 ^b (2,250-6,534)	3,454 ^b (1,504-7,144)
Monocytes (x10 ⁹ /L)	0.725 ^a (0.172-2256)	0.3585 ^b (0.138-0.830)	0.334 ^b (0.86-0.735)
Hb Fetal (%)	10.2 ^a (3.8-24.2)	0.0(0.0-3.0)	2.15 ^c (0.0-2.6)
Hb S (%)	83.5 ^a (55.3-93.9)	36.6 ^b (28.7-40.5)	0.0
Hb A (%)	0.0(0-33)	58.9 ^a (55-65.5)	95.4 ^b (95-97.6)

Hb - hemoglobin, MCH -mean corpuscular hemoglobin, MCHC -mean corpuscular hemoglobin concentration, MCV -mean corpuscular volume, Max - maximum, Med - median, Min - minimum, TBARS - thiobarbituric acid reactive species, TEAC - Trolox equivalent antioxidant capacity. The same superscript letters indicate non-significant differences ($p > 0.05$) while different superscript letters indicate significant differences ($p < 0.05$). Kruskal-Wallis and Mann Whitney tests with a 0.05 significance level.

cyte levels ($p = 0.03$) (Table 4). When grouped together, the Bantu and Benin haplotypes showed significant increases in LPO and TEAC levels, and a significant reduction in melatonin levels ($p < 0.05$) as compared to the control group (Hb AA) (Table 5). There were no significant differences in the clinical characteristics of the Bantu/Bantu and

Bantu/Benin groups, or between HbSS/ $\alpha^{3.7kb}$ patients and those without thalassemia ($p > 0.05$) (Table 6).

Spearman correlation analyses revealed positive correlations between the TBARS and HCM values in the Bantu/Bantu group ($r = 0.84$; $p = 0.03$). There were also negative correlations between TEAC and melatonin levels

Table 4 - Redox profile and hematological characteristics of SCA patients according to β^S gene haplotypes, represented by median values (minimum and maximum).

Features	Bantu/Bantu (N = 6)	Bantu/Benin (N = 7)	p
TBARS (ng/mL)	910.9(496.8-1970)	842.3(635.5-2,203)	NS
TEAC (mM)	2.0(2.0-2.1)	1.9(1.9-2.1)	NS
Melatonin (pg/mL)	44.8(17-164)	39.2(9.63-74.1)	NS
Hb (g/dL)	8.1(6.4-9.7)	7.6(5.9-9.0)	NS
MCV (fL)	102.7(89.1-118.6)	92.9(79.5-121.9)	NS
MCH (pg)	34.4(28.2-38.3)	30.2(24.8-40.4)	NS
MCHC (%)	32.4(31-35.6)	33.2(31.2-34.7)	NS
Reticulocytes (%)	13.4(7.3-36.9)	11.2(6.3-28.6)	NS
Hb F (%)	9.5(3.8-15.9)	14.3(1.2-24.2)	NS
Hb S (%)	85.1(55.3-89.0)	81.7(71.7-93.9)	NS
Leukocytes (x10 ⁹ /L)	12,150(8,800-18,800)	13,100(8,600-15,500)	NS
Neutrophils (x10 ⁹ /L)	6,894(4,536-11,092)	5,633(4,386-8,370)	NS
Monocytes (x10 ⁹ /L)	1,202(0.54-2,256)	0.655(0.172-1,085)	0.03

Hb - hemoglobin, MCH - mean corpuscular hemoglobin, MCHC - mean corpuscular hemoglobin concentration, MCV - mean corpuscular volume, SCA - sickle cell anemia, TBARS - thiobarbituric acid reactive species, TEAC - Trolox equivalent antioxidant capacity. NS - non-significant ($p > 0.05$). Mann Whitney test with a 0.05 significance level.

Table 5 - Antioxidant capacity and lipid peroxidation in Bantu and Benin haplotypes as compared to the control group (Hb AA).

	BAN/BEN+BAN/BAN (n = 13) Med (min-max)	Hb AA (n = 30) Med (min-max)	p
TEAC (mM)	2.0 (1.9-2.1)	1.9 (1.7-2.0)	0.002
Melatonin (pg/mL)	42.6 (9.6-164)	146.5 (39.9-671.6)	0.007
TBARS (ng/mL)	842.3 (496.8-2203)	485.0 (270.2-676.3)	< 0.001

BAN/BAN -Bantu/Bantu, BAN/BEN -Bantu/Benin, Max - maximum, Med - median, Min - minimum, TBARS - thiobarbituric acid reactive species, TEAC - Trolox equivalent antioxidant capacity. Mann Whitney test with a 0.05 significance level.

Table 6 - Hydroxyurea (HU) use and frequency of clinical manifestations in SCA patients according to β^S gene haplotype, $-\alpha^{3.7kb}$ thalassemia coinheritance and normal α genotype.

	BAN/BAN (n = 6) (%)	BAN/BEN (n = 7) (%)	p	SS/ $-\alpha^{3.7}$ (n = 3) (%)	SS/($\alpha\alpha/\alpha\alpha$) (n = 14) (%)	p
Leg ulcers	0	14.3	-	33.3	0	-
Pneumonia	16.6	14.3	NS	33.3	25	NS
Splenectomy	16.6	28.5	NS	33.3	28.5	NS
Cholecystectomy/ cholelithiasis	16.6	42.8	NS	33.3	28.5	NS
Pleural effusion	16.6	0	-	0	7.1	-
Acute thoracic syndrome	0	0	-	0	7.1	-
Cardiac complications	33.3	57.1	NS	66.7	35.7	NS
Osteonecrosis	33.3	14.3	NS	0	28.5	-
Osteomyelitis	0	16.6	-	0	7.1	-
Joint/muscle pain	50	57.1	NS	66.7	50	NS
Urinary infection	16.6	0	-	0	7.1	-
HU use	66.6	28.6	NS	0	64.2	NS

BAN/BAN - Bantu/Bantu, BAN/BEN - Bantu/Benin, NS -non-significant ($p > 0.05$). Fisher's Exact test with a significance of $p < 0.05$.

($r = -0.91$; $p = 0.01$), and between Hb F and reticulocytes ($r = -0.9$; $p < 0.001$) in this group. A strong correlation trending towards significance was also found between monocytes and reticulocytes ($r = 0.77$; $p = 0.07$). In the Bantu/Benin group, there were positive correlations between TEAC and reticulocytes ($r = 0.73$; $p = 0.05$), TBARS and reticulocytes ($r = 0.82$; $p = 0.02$), TBARS and monocytes ($r = 0.73$; $p = 0.05$) and Hb F and total Hb ($r = 0.83$; $p = 0.01$). The data suggested that LPO was directly associated with the degree of hemolysis. In the SS/ $-\alpha^{3.7}$ group, there were positive correlations between TBARS and TEAC ($r = 0.9$; $p < 0.001$) and between TBARS and reticulocytes ($r = 0.9$; $p < 0.001$), and negative correlations between TBARS and total Hb ($r = -0.9$; $p < 0.001$). In the Hb SS/($\alpha\alpha/\alpha\alpha$) group, there was a negative correlation between Hb F and Hb S ($r = -0.64$; $p = 0.012$).

Discussion

This is the first report on haplotype frequency in Hb SS individuals in the state of Paraná. The results indicate that the Bantu haplotype, followed by the Benin haplotype, was the most frequent in the northwest region of Paraná. The most frequent genotype was Bantu/Benin. The haplo-

type frequency identified here was similar to that reported for the states of São Paulo (Figueiredo *et al.*, 1996) and Rio de Janeiro (Silva Filho *et al.*, 2010). For many years, slaves arrived in Brazil on forced migration routes between the Benin bay in Ghana and Nigeria to the northeastern region of Brazil, and from Congo and Angola to Rio de Janeiro (Silva Filho *et al.*, 2010). Rio de Janeiro was a major slave distribution center, sending slaves especially to the states of São Paulo and Rio Grande do Sul. The high frequency of Bantu followed by Benin haplotypes identified in this study agrees with historical records of the slave trade and internal migration between states in Brazil.

The present findings also agree with the scientific literature, as other studies have reported similar results for the association between the Bantu/Bantu and Bantu/Benin haplotypes, increased LPO and reduced antioxidant defense (Rusanova *et al.*, 2010). The Bantu haplotype is associated with a lower concentration of Hb F and greater clinical severity than the Benin haplotype (Powars, 1991). However, our data showed no significant differences in the Hb F values, global redox status or clinical manifestations between homozygous Bantu haplotypes and heterozygous Bantu/Benin haplotypes. These results corroborate data from other studies (Rieder *et al.*, 1991; Figueiredo *et al.*,

1996; Silva and Gonçalves, 2010). Based on our results, it is possible that neither the Bantu/Benin nor the Bantu/Bantu haplotypes protect against oxidative damage and clinical features of SCA. If the Senegal haplotype had been present in the sample, it would have been possible to draw more accurate conclusions regarding the effect of these polymorphisms on oxidative and clinical-laboratorial characteristics.

The significantly reduced serum melatonin concentration in patients with SCA corroborates a previous study by Shimauti *et al.* (2010) that also found an association between low melatonin concentrations and this illness. The decrease in serum melatonin together with the elevated TEAC and LPO levels observed here agree with results from other investigations (Gitto *et al.*, 2001; Reiter *et al.*, 2005). Although serum melatonin levels tend to decrease as age increases (Waldhauser *et al.*, 1988; Zhdanova *et al.*, 1998), melatonin concentrations in individuals with SCA (Hb SS) have been found to be equally low in all age-groups (Shimauti *et al.*, 2010). Studies have shown that, in conditions of intense oxidative stress, interactions with ROS lead to the metabolization of melatonin and subsequent reductions in its serum concentration (Tan *et al.*, 2007). Melatonin also increases the total antioxidant capacity, possibly by acting in synergy with classic antioxidants.

In SCA, monocytes are responsible for the activation of endothelial cells, and act as a trigger for the translocation of the nuclear transcription factor NF- κ B (Belcher *et al.*, 2000). The activated vascular endothelium expresses the adhesion molecules ICAM-I, VCAM-I, P-selectin and E-selectin, and the procoagulant tissue factor, all which play an important role in the physiopathology of vascular occlusion in Hb SS individuals. The strong positive correlation between monocytes and reticulocytes in the Bantu/Bantu group, and between TBARS, reticulocytes and monocytes in the Bantu/Benin group suggests that higher hemolytic rates are associated with increased LPO and inflammatory responses. These data suggest a vulnerability of these genotypes to vasculopathy.

The frequency of coinheritance of Hb S with $-\alpha^{3.7\text{kb}}$ observed here was similar to that reported for the states of São Paulo and Rio de Janeiro (Figueiredo *et al.*, 1996; Silva Filho *et al.*, 2010). In these individuals, the most commonly observed haplotype was Bantu/Benin, as reported in studies done in the city of Salvador, in the state of Bahia, and in the state of São Paulo (Lyra *et al.*, 2005). Investigations of the interaction between Hb SS and alpha thalassemia ($-\alpha^{3.7\text{kb}}$) and its effect on clinical and hematological factors have produced conflicting results. Some studies have reported reduced hemolytic rates, increased total Hb concentration and a protective effect against clinical manifestations of the disease (Higgs *et al.*, 1982; Powars, 1991; Belisario *et al.*, 2010; Fertrin and Costa, 2010), while other studies have found no differences between individuals with and without the $-\alpha^{3.7}$ mutation ($\alpha\alpha/\alpha\alpha$) (Mouele *et al.*, 1999).

Previous studies have reported on the importance of alpha thalassemia in the modulation of SCA (Steinberg, 2005; Belisario *et al.*, 2010); however, our study failed to demonstrate significant differences in the frequency of clinical complications between groups with and without $-\alpha^{3.7}$ thalassemia coinheritance, possibly because of our small sample size. Nonetheless, the inverse correlation detected between TBARS and total Hb and the positive correlation between reticulocytes and TBARS suggest that $-\alpha^{3.7}$ thalassemia may decrease hemolysis and attenuate lipid peroxidation. It is also interesting to note that, in the present sample, the use of HU, a chemotherapeutic agent used to increase the Hb F concentration and attenuate hemolysis, was more common in SS individuals without the $-\alpha^{3.7\text{kb}}$ mutation and in patients with the Bantu/Bantu genotype, which is generally associated with a worse clinical outcome.

The ability of HU to increase Hb F levels varies among patients and the duration of treatment with this chemotherapeutic agent influences the increase in Hb F. In order to achieve the expected benefits, the treatment should last for at least two years (Davies and Gilmore, 2003). As shown here, most patients (66.6%) with Hb SS who were receiving HU had been under treatment for less than two years. Therefore, treatment duration may be one of the determinant factors for the lack of significant differences in Hb F levels between patients receiving HU or not.

In conclusion, the results of this study provide a relevant contribution to our understanding of the pattern of colonization and to the anthropological and historical background of the Afro-Brazilian population in the state of Paraná. Our data suggest that the antioxidant defense in SCA patients with the Bantu/Bantu or Bantu/Benin genotype is insufficient to control oxidative stress. Melatonin, a potent antioxidant and anti-inflammatory agent, is a possible treatment option for SCA patients with low serum melatonin levels. However, the protective effect of melatonin against oxidative stress in individuals with sickle cell disease requires further study. More comprehensive investigations using data from other regions in the state of Paraná must be done to better characterize the frequency of β^S gene haplotypes and of $-\alpha^{3.7\text{kb}}$ thalassemia coinheritance, as well as their effects on oxidative stress and phenotypic expression in patients with Hb SS.

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